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Nucleotide Sequences pXP10 (SEQ ID No: 111)

1	GACGAAAGGG	CCTCGTGATA	CGCCTATTTT	TATAGGTAA	TGTCATGATA
	CTGCTTTCCC	GGAGCACTAT	GCGGATAAAA	ATATCCAATT	ACAGTACTAT
51	ATAATGGTTT	CTTAGACGTC	AGGTGGCACT	TTTCGGGGAA	ATGTGCGCGG
	TATTACCAAA	GAATCTGCAG	TCCACCGTGA	AAAGCCCCTT	TACACGCGCC
101	AACCCCTATT	TGTTTATTTT	TCTAAATACA	TTCAAATATG	TATCCGCTCA
	TTGGGGATAA	ACAAATAAAA	AGATTTATGT	AAGTTTATAC	ATAGGCGAGT
151	TGAGACAATA	ACCCTGATAA	ATGCTTCAAT	AATATTGAAA	AAGGAAGAGT
	ACTCTGTTAT	TGGGACTATT	TACGAAGTTA	TTATAACTTT	TTCTTTCTCA
201	ATGAGTATTC	AACATTTCCG	TGTCGCCCTT	ATTCCCTTTT	TTGCGGCATT
	TACTCATAAG	TTGTAAAGGC	ACAGCGGGAA	TAAGGGAAAA	AACGCCGTAA
251	TTGCCTTCCT	GTTTTTGCTC	ACCCAGAAAC	GCTGGTGAAA	GTAAAAGATG
	AACGGAAGGA	CAAAAACGAG	TGGGTCTTTG	CGACCACTTT	CATTTTCTAC
301	CTGAAGATCA	GTTGGGTGCT	CGAGTGGGTT	ACATCGAACT	GGATCTCAAC
	GACTTCTAGT	CAACCCACGA	GCTCACCCAA	TGTAGCTTGA	CCTAGAGTTG
351	AGCGGTAAGA	TCCTTGAGAG	TTTTCGCCCC	GAAGAACGTT	TTCCAATGAT
	TCGCCATTCT	AGGAACTCTC	AAAAGCGGGG	CTTCTTGCAA	AAGGTTACTA
401	GAGCACTTTT	AAAGTTCTGC	TATGTGGCGC	GGTATTATCC	CGTATTGACG
	CTCGTGAAAA	TTTCAAGACG	ATACACCGCG	CCATAATAGG	GCATAACTGC
451	CCGGGCAAGA	GCAACTCGGT	CGCCGCATAC	ACTATTCTCA	GAATGACTTG
	GGCCCGTTCT	CGTTGAGCCA	GCGGCGTATG	TGATAAGAGT	CTTACTGAAC
501	GTTGAGTACT	CACCAGTCAC	AGAAAAGCAT	CTTACGGATG	GCATGACAGT
	CAACTCATGA	GTGGTCAGTG	TCTTTTCGTA	GAATGCCTAC	CGTACTGTCA
551	AAGAGAATTA	TGCAGTGCTG	CCATAACCAT	GAGTGATAAC	ACTGCGGCCA
	TTCTCTTAAT	ACGTCACGAC	GGTATTGGTA	CTCACTATTG	TGACGCCGGT
601	ACTTACTTCT	GACAACGATC	GGAGGACCGA	AGGAGCTAAC	CGCTTTTTTG
	TGAATGAAGA	CTGTTGCTAG	CCTCCTGGCT	TCCTCGATTG	GCGAAAAAAC
651	CACAACATGG	GGGATCATGT	AACTCGCCTT	GATCGTTGGG	AACCGGAGCT
	GTGTTGTACC	CCCTAGTACA	TTGAGCGGAA	CTAGCAACCC	TTGGCCTCGA
701	GAATGAAGCC	ATACCAAACG	ACGAGCGTGA	CACCACGATG	CCTGTAGCAA
	CTTACTTCGG	TATGGTTTGC	TGCTCGCACT	GTGGTGCTAC	GGACATCGTT
751	TGGCAACAAC	GTTGCGCAAA	CTATTAAGTG	GCGAACTACT	TACTCTAGCT
	ACCGTTGTTG	CAACGCGTTT	GATAATTGAC	CGCTTGATGA	ATGAGATCGA
801	TCCCGGCAAC	AATTAATAGA	CTGGATGGAG	GCGGATAAAG	TTGCAGGACC
	AGGGCCGTTG	TTAATTATCT	GACCTACCTC	CGCCTATTTC	AACGTCCTGG
851	ACTTCTGCGC	TCGGCCCTTC	CGGCTGGCTG	GTTTATTGCT	GATAAATCTG
	TGAAGACGCG	AGCCGGGAAG	GCCGACCGAC	CAAATAACGA	CTATTTAGAC
901	GAGCCGGTGA	GCGTGGGTCT	CGCGGTATCA	TTGCAGCACT	GGGGCCAGAT
	CTCGGCCACT	CGCACCCAGA	GCGCCATAGT	AACGTCGTGA	CCCCGGTCTA
951	GGTAAGCCCT	CCCGTATCGT	AGTTATCTAC	ACGACGGGGA	GTCAGGCAAC
	CCATTCGGGA	GGGCATAGCA	TCAATAGATG	TGCTGCCCCT	CAGTCCGTTG
1001	TATGGATGAA	CGAAATAGAC	AGATCGCTGA	GATAGGTGCC	TCACTGATTA
	ATACCTACTT	GCTTTATCTG	TCTAGCGACT	CTATCCACGG	AGTGACTAAT
1051	AGCATTGGTA	ACTGTCAGAC	CAAGTTTACT	CATATATACT	TTAGATTGAT
	TCGTAACCAT	TGACAGTCTG	GTTCAAATGA	GTATATATGA	AATCTAACTA
1101	TTAAACTTTC	ATTTTTAATT	TAAAAGGATC	TAGGTGAAGA	TCCTTTTTGA
	AATTTTGAAG	TAAAAATTAA	ATTTTCCTAG	ATCCACTTCT	AGGAAAAACT
1151	TAATCTCATG	ACCAAAATCC	CTTAACGTGA	GTTTTCGTTC	CACTGAGCGT
	ATTAGAGTAC	TGGTTTTAGG	GAATTGCACT	CAAAAGCAAG	GTGACTCGCA
1201	CAGACCCCGT	AGAAAAGATC	AAAGGATCTT	CTTGAGATCC	TTTTTTTCTG
	GTCTGGGGCA	TCTTTTCTAG	TTTCCTAGAA	GAACCTAGAG	AAAAAAGAC
1251	CGCGTAATCT	GCTGCTTGCA	AACAAAAAAA	CCACCCTAGC	CAGCGGTGGT
	GCGCATTAGA	CGACGAACGT	TTGTTTTTTT	GGTGGCGATG	GTCGCCACCA
1301	TTGTTTGCCG	GATCAAGAGC	TACCAACTCT	TTTTCCGAAG	GTAAGTGGCT
	AACAAACGGC	CTAGTTCTCG	ATGGTTGAGA	AAAAGGCTTC	CATTGACCGA
1351	TCAGCAGAGC	GCAGATACCA	AATACTGTCC	TTCTAGTGTA	GCCGTAGTTA

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1401	AGTCGTCTCG	CGTCTATGGT	TTATGACAGG	AAGATCACAT	CGGCATCAAT
	GGCCACCACT	TCAAGAACTC	TGTAGCACCG	CCTACATACC	TCGCTCTGCT
	CCGGTGGTGA	AGTTCTTGAG	ACATCGTGGC	GGATGTATGG	AGCGAGACGA
1451	AATCCTGTTA	CCAGTGGCTG	CTGCCAGTGG	CGATAAGTCG	TGTCTTACCG
	TTAGGACAAT	GGTCACCGAC	GACGGTCACC	GCTATTCAGC	ACAGAATGGC
1501	GGTTGGACTC	AAGACGATAG	TTACCGGATA	AGGCGCAGCG	GTCGGGCTGA
	CCAACCTGAG	TTCTGCTATC	AATGGCCTAT	TCCGCGTCGC	CAGCCCGACT
1551	ACGGGGGGTT	CGTGCATACA	GCCCAGCTTG	GAGCGAACGA	CCTACACCGA
	TGCCCCCAA	GCACGTATGT	CGGGTCGAAC	CTCGCTTGCT	GGATGTGGCT
1601	ACTGAGATAC	CTACAGCGTG	AGCTATGAGA	AAGCGCCACG	CTTCCCGAAG
	TGACTCTATG	GATGTCGCAC	TCGATACTCT	TTCGCGGTGC	GAAGGGCTTC
1651	GGAGAAAGGC	GGACAGGTAT	CCGGTAAGCG	GCAGGGTCGG	AACAGGAGAG
	CCTCTTTCCG	CCTGTCCATA	GGCCATTTCG	CGTCCCAGCC	TTGTCCTCTC
1701	CGCACGAGGG	AGCTTCCAGG	GGGAAACGCC	TGGTATCTTT	ATAGTCCTGT
	GCGTGCTCCC	TCGAAGGTCC	CCCTTTGCGG	ACCATAGAAA	TATCAGGACA
1751	CGGGTTTTCG	CACCTCTGAC	TTGAGCGTCG	ATTTTTGTGA	TGCTCGTCAG
	GCCCAAAGCG	GTGGAGACTG	AACTCGCAGC	TAAAAACACT	ACGAGCAGTC
1801	GGGGGCGGAG	CCTATGGAAA	AACGCCAGCA	ACGCGGCCTT	TTTACGGTTC
	CCCCCGCCTC	GGATACCTTT	TTGCGGTCTG	TGCGCCGGAA	AAATGCCAAG
1851	CTGGCCTTTT	GCTGGCCTTT	TGCTCACATG	TTCTTTCTTG	CGTTATCCCC
	GACCGGAAAA	CGACCGGAAA	ACGAGTGTAC	AAGAAAGGAC	GCAATAGGGG
1901	TGATTCTGTG	GATAACCGTA	TTACCGCCTT	TGAGTGAGCT	GATACCGCTC
	ACTAAGACAC	CTATTGGCAT	AATGGCGGAA	ACTCACTCGA	CTATGGCGAG
1951	GCCGCAGCCG	AACGACCGAG	CGCAGCGAGT	CAGTGAGCGA	GGAAGCGGAA
	CGGCGTCGGC	TTGCTGGCTC	GCGTCGCTCA	GTCACTCGCT	CCTTCGCCTT
2001	GAGCGCCCAA	TACGCAAACC	GCCTCTCCCC	GCGCGTTGGC	CGATTCAATTA
	CTCGCGGGTT	ATGCGTTTGG	CGGAGAGGGG	CGCGCAACCG	GCTAAGTAAT
2051	ATGCAGCTGG	CACGACAGGT	TTCCCGACTG	GAAAGCGGGC	AGTGAGCGCA
	TACGTCAAGC	GTGCTGTCCA	AAGGGCTGAC	CTTTTCGCCC	TCACTCGCGT
2101	ACGCAATTAA	TGTGAGTTAG	CTCACTCATT	AGGCACCCCA	GGCTTTACAC
	TGCGTTAATT	ACACTCAATC	GAGTGAGTAA	TCCGTGGGGT	CCGAAATGTG
2151	TTTATGCTTC	CGGCTCGTAT	GTTGTGTGGA	ATTGTGAGCG	GATAACAATT
	AAATACGAAG	GCCGAGCATA	CAACACACCT	TAACACTCGC	CTATTGTTAA
2201	TCACACAGGA	AACAGCTATG	ACCATGATTA	CGCCAAGCTT	TGGAGCCTTT
	AGTGTGTCTT	TTGTCGATAC	TGGTACTAAT	GCGGTTTCGA	ACCTCGGAAA
2251	TTTTTGGAGA	TTTTCAACGT	GAAAAAATTA	TTATTTCGAA	TTCTTTTAGT
	AAAAACCTCT	AAAAGTTGCA	CTTTTTTAAT	AATAAGCGTT	AAGGAAATCA
2301	TGTTTCCTTT	TATGCGGCCC	AGCCGGCCAT	GGCCCAGGTC	CAGTCGACAG
	ACAAGGAAAG	ATACGCCGGG	TCGGCCGGTA	CCGGGTCCAG	GTCAGCTGTC
2351	GTGGAGGCGG	TTCAGGCGGA	GGTGGCTCTG	GCGGTGGCGG	AAGTGCCTCT
	CACCTCCGCC	AAGTCCGCCT	CCACCGAGAC	CGCCACCGCC	TTCACGTGAG
2401	ATCAAACGGC	GGCCGCAGGT	GCGCCGGTGC	CGTATCCGGA	TCCGCTGGAA
	TAGTTTGCCG	CCGGCGTCCA	CGCGGCCACG	GCATAGGCCT	AGGCGACCTT
2451	CCGCGTGCCG	CATAGGCTGG	CGGCGGCTCT	GGTGGTGGTT	CTGGTGGCGG
	GGCGCACGGC	GTATCCGACC	GCCGCCGAGA	CCACCACCAA	GACCACCGCC
2501	CTCTGAGGGT	GGCGGCTCTG	AGGGTGGCGG	TTCTGAGGGT	GGCGGCTCTG
	GAGACTCCCA	CCGCCGAGAC	TCCCACCGCC	AAGACTCCCA	CCGCCGAGAC
2551	AGGGTGGCGG	TTCCGGTGGC	GGCTCCGGTT	CCGGTGATTT	TGATTATGAA
	TCCCACCGCC	AAGGCCACCG	CCGAGGCCAA	GGCCACTAAA	ACTAATACTT
2601	AAAATGGCAA	ACGCTAATAA	GGGGGCTATG	ACCGAAAATG	CCGATGAAAA
	TTTTACCGTT	TGCGATTATT	CCCCCGATAC	TGGCTTTTAC	GGCTACTTTT
2651	CGCGCTACAG	TCTGACGCTA	AAGGCAAAC	TGATTCTGTC	GCTACTGATT
	GCGCGATGTC	AGACTGCGAT	TTCCGTTTGA	ACTAAGACAG	CGATGACTAA
2701	ACGGTGCTGC	TATCGATGGT	TTCAATTGGT	ACGTTTCCGG	CCTTGCTAAT
	TGCCACGACG	ATAGCTACCA	AAGTAACCAC	TGCAAAGGCC	GGAACGATTA
2751	GGTAATGGTG	CTACTGGTGA	TTTTGCTGGC	TCTAATTCCC	AAATGGCTCA
	CCATTACCAC	GATGACCACT	AAAACGACCG	AGATTAAAGG	TTTACCGAGT

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2801	AGTCGGTGAC	GGTGATAATT	CACCTTTAAT	GAATAATTTT	CGTCAATATT
	TCAGCCACTG	CCACTATTAA	GTGGAAATTA	CTTATTAAAG	GCAGTTATAA
2851	TACCTTCTTT	GCCTCAGTCG	GTTGAATGTC	GCCCTTATGT	CTTTGGCGCT
	ATGGAAGAAA	CGGAGTCAGC	CAACTTACAG	CGGGAATACA	GAAACCGCGA
2901	GGTAAACCAT	ATGAATTTTC	TATTGATTGT	GACAAAATAA	ACTTATTCCG
	CCATTTGGTA	TACTTAAAAG	ATAACTAACA	CTGTTTTATT	TGAATAAGGC
2951	TGGTGTCTTT	GCGTTTCTTT	TATATGTTGC	CACCTTTATG	TATGTATTTT
	ACCACAGAAA	CGCAAAGAAA	ATATACAACG	GTGGAAATAC	ATACATAAAA
3001	CGACGTTTGC	TAACATACTG	CGTAATAAGG	AGTCTTAATA	AGAATTCACT
	GCTGCAAACG	ATTGTATGAC	GCATTATTCC	TCAGAATTAT	TCTTAAGTGA
3051	GGCGCTCGTT	TTACAACGTC	GTGACTGGGA	AAACCCTGGC	GTTACCCAAC
	CCGGCAGCAA	AATGTTGCAG	CAGTACCCTT	TTTGGGACCG	CAATGGGTTG
3101	TTAATCGCCT	TGCAGCACAT	CCCCCTTTTC	CCAGCTGGCG	TAATAGCGAA
	AATTAGCGGA	ACGTCGTGTA	GGGGGAAAGC	GGTCGACCGC	ATTATCGCTT
3151	GAGGCCGCA	CCGATCGCCC	TTCCCAACAG	TTGCGCAGCC	TGAATGGCGA
	CTCCGGGCGT	GGCTAGCGGG	AAGGGTTGTC	AACGCGTCGG	ACTTACCGCT
3201	ATGGCGCCTG	ATGCGGTATT	TTCTCCTTAC	GCATCTGTGC	GGTATTTTAC
	TACCGCGGAC	TACGCCATAA	AAGAGGAATG	CGTAGACACG	CCATAAAGTG
3251	ACCGCATACG	TCAAAGCAAC	CATAGTACGC	GCCCTGTAGC	GGCGCATTAA
	TGGCGTATGC	AGTTTCGTTG	GTATCATGCG	CGGGACATCG	CCGCGTAATT
3301	GCCCGGCGGG	TGTGGTGGTT	ACGCGCAGCG	TGACCGCTAC	ACTTGCCAGC
	CGGGCCGCCC	ACACCACCAA	TGCGCGTCGC	ACTGGCGATG	TGAACGGTCG
3351	GCCCTAGCCC	CCGCTCCTTT	CGCTTTCTTC	CCTTCCTTTC	TCGCCACGTT
	CGGGATCGGG	GGCGAGGAAA	GCGAAAGAAG	GGAAGGAAAG	AGCGGTGCAA
3401	CGCCGGCTTT	CCCCGTCAAG	CTCTAAATCG	GGGGCTCCCT	TTAGGGTTCC
	GCGGCCGAAA	GGGGCAGTTC	GAGATTTAGC	CCCCGAGGGA	AATCCCAAGG
3451	GATTTAGTGC	TTTACGGCAC	CTCGACCCCA	AAAAACTTGA	TTTGGGTGAT
	CTAAATCACG	AAATGCCGTG	GAGCTGGGGT	TTTTTGAACT	AAACCCACTA
3501	GGTTCACGTA	GTGGGCCATC	GCCCTGATAG	ACGGTTTTTC	GTCCTTTGAC
	CCAATCGCAT	CACCCGGTAG	CGGGACTATC	TGCCAAAAAG	CAGGAAACTG
3551	GTTTCGAGTCC	ACGTTCTTTA	ATAGTGGACT	CTTGTTCCAA	ACTGGAACAA
	CAAGCTCAGG	TGCAAGAAAT	TATCACCTGA	GAACAAGGTT	TGACCTTGTT
3601	TACTCAACCC	TATCTCGGGC	TATTCTTTTG	ATTTATAAGG	GATTTTGCCG
	ATGAGTTGGG	ATAGAGCCCG	ATAAGAAAAC	TAAATATTCC	CTAAAACGGC
3651	ATTTTCGGCT	ATTGGTTAAA	AAATGAGCTG	ATTTAACAAA	AATTTAACGC
	TAAAGCCGGA	TAACCAATTT	TTTACTCGAC	TAAATTGTTT	TTAAATTGCG
3701	GAATTTTAAAC	AAAATATTAA	CGTTTACAAT	TTTATGGTGC	AGTCTCAGTA
	CTTAAATTTG	TTTTATAATT	GCAAATGTTA	AAATACCACG	TCAGAGTCAT
3751	CAATCTGCTC	TGATGCCGCA	TAGTTAAGCC	AGCCCCGACA	CCCGCCAACA
	GTTAGACGAG	ACTACGGCGT	ATCAATTCGG	TCGGGGCTGT	GGGCGGTTGT
3801	CCCGCTGACG	CGCCCTGACG	GGCTTGTCTG	CTCCCGGCAT	CCGCTTACAG
	GGGCGACTGC	GCGGGACTGC	CCGAACAGAC	GAGGGCCGTA	GGCGAATGTC
3851	ACAAGCTGTG	ACCGTCTCCG	GGAGCTGCAT	GTGTCAGAGG	TTTTCACCGT
	TGTTTCGACAC	TGGCAGAGGC	CCTCGACGTA	CACAGTCTCC	AAAAGTGGCA
3901	CATCACCGBA	ACGCGCGA			
	GTAGTGGCTT	TGCGCGCT			

Fig. 7b

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Nucleotide Sequences pXP14 (SEQ ID No: 112)

1	GACGAAAGGG	CCTCGTGATA	CGCCTATTTT	TATAGGTAA	TGTCATGATA
	CTGCTTTCCC	GGAGCACTAT	GCGGATAAAA	ATATCCAATT	ACAGTACTAT
51	ATAATGGTTT	CTTAGACGTC	AGGTGGCACT	TTTCGGGGAA	ATGTGCGCGG
	TATTACCAAA	GAATCTGCAG	TCCACCGTGA	AAAGCCCCTT	TACACGCGCC
101	AACCCCTATT	TGTTTATTTT	TCTAAATACA	TTCAAATATG	TATCCGCTCA
	TTGGGGATAA	ACAAATAAAA	AGATTTATGT	AAGTTTATAC	ATAGGCGAGT
151	TGAGACAATA	ACCCTGATAA	ATGCTTCAAT	AATATTGAAA	AAGGAAGAGT
	ACTCTGTTAT	TGGGACTATT	TACGAAGTTA	TTATAACTTT	TTCCTTCTCA
201	ATGAGTATTC	AACATTTCCG	TGTCGCCCTT	ATTCCCTTTT	TTGCGGCATT
	TACTCATAAG	TTGTAAAGGC	ACAGCGGGAA	TAAGGGAAAA	AACGCCGTAA
251	TTGCCTTCCT	GTTTTTGCTC	ACCCAGAAAC	GCTGGTGAAA	GTAAAAGATG
	AACGGAAGGA	CAAAAACGAG	TGGGTCTTTG	CGACCACTTT	CATTTTCTAC
301	CTGAAGATCA	GTTGGGTGCT	CGAGTGGGTT	ACATCGAACT	GGATCTCAAC
	GACTTCTAGT	CAACCCACGA	GCTCACCCAA	TGTAGCTTGA	CCTAGAGTTG
351	AGCGGTAAAG	TCCTTGAGAG	TTTTCGCCCC	GAAGAACGTT	TTCCAATGAT
	TCGCCATTCT	AGGAACTCTC	AAAAGCGGGG	CTTCTTGCAA	AAGGTTACTA
401	GAGCACTTTT	AAAGTTCTGC	TATGTGGCGC	GGTATTATCC	CGTATTGACG
	CTCGTGAAAA	TTTCAAGACG	ATACACCGCG	CCATAATAGG	GCATAACTGC
451	CCGGGCAAGA	GCAACTCGGT	CGCCGCATAC	ACTATTCTCA	GAATGACTTG
	GGCCCGTTCT	CGTTGAGCCA	GCGGCGTATG	TGATAAGAGT	CTTACTGAAC
501	GTTGAGTACT	CACCAGTCAC	AGAAAAGCAT	CTTACGGATG	GCATGACAGT
	CAACTCATGA	GTGGTCAGTG	TCTTTTCGTA	GAATGCCTAC	CGTACTGTCA
551	AAGAGAATTA	TGCAGTGCTG	CCATAACCAT	GAGTGATAAC	ACTGCGGCCA
	TTCTCTTAAT	ACGTCACGAC	GGTATTGGTA	CTCACTATTG	TGACGCCGGT
601	ACTTACTTCT	GACAACGATC	GGAGGACCGA	AGGAGCTAAC	CGCTTTTTTG
	TGAATGAAGA	CTGTTGCTAG	CCTCCTGGCT	TCCTCGATTG	GCGAAAAAAC
651	CACAACATGG	GGGATCATGT	AACTCGCCTT	GATCGTTGGG	AACCGGAGCT
	GTGTTGTACC	CCCTAGTACA	TTGAGCGGAA	CTAGCAACCC	TTGGCCTCGA
701	GAATGAAGCC	ATACCAAACG	ACGAGCGTGA	CACCACGATG	CCTGTAGCAA
	CTTACTTCGG	TATGGTTTGC	TGCTCGCACT	GTGGTGCTAC	GGACATCGTT
751	TGGCAACAAC	GTTGCGCAAA	CTATTAAC TG	GCGAACTACT	TACTCTAGCT
	ACCGTTGTTG	CAACGCGTTT	GATAATTGAC	CGCTTGATGA	ATGAGATCGA
801	TCCCGGCAAC	AATTAATAGA	CTGGATGGAG	GCGGATAAAG	TTGCAGGACC
	AGGGCCGTTG	TTAATTATCT	GACCTACCTC	CGCCTATTTT	AACGTCCTGG
851	ACTTCTGCGC	TCGGCCCTTC	CGGCTGGCTG	GTTTATTGCT	GATAAATCTG
	TGAAGACGCG	AGCCGGGAAG	GCCGACCGAC	CAAATAACGA	CTATTTAGAC
901	GAGCCGGTGA	GCGTGGGTCT	CGCGGTATCA	TTGCAGCACT	GGGGCCAGAT
	CTCGGCCACT	CGCACCCAGA	GCGCCATAGT	AACGTCGTGA	CCCCGGTCTA
951	GGTAAGCCCT	CCCGTATCGT	AGTTATCTAC	ACGACGGGGA	GTCAGGCAAC
	CCATTCGGGA	GGGCATAGCA	TCAATAGATG	TGCTGCCCTT	CAGTCCGTTG
1001	TATGGATGAA	CGAAATAGAC	AGATCGCTGA	GATAGGTGCC	TCACTGATTA
	ATACCTACTT	GCTTTATCTG	TCTAGCGACT	CTATCCACGG	AGTGACTAAT
1051	AGCATTGGTA	ACTGTCAGAC	CAAGTTTACT	CATATATACT	TTAGATTGAT
	TCGTAACCAT	TGACAGTCTG	GTTCAAATGA	GTATATATGA	AATCTAACTA
1101	TTAAAACCTT	ATTTTAAATT	TAAAAGGATC	TAGGTGAAGA	TCCTTTTTGA
	AATTTTGAAG	TAAAAATTAA	ATTTTCCTAG	ATCCACTTCT	AGGAAAAACT
1151	TAATCTCATG	ACCAAATCC	CTTAACGTGA	GTTTTCGTTC	CACTGAGCGT
	ATTAGAGTAC	TGGTTTTAGG	GAATTGCACT	CAAAAGCAAG	GTGACTCGCA
1201	CAGACCCCGT	AGAAAAGATC	AAAGGATCTT	CTTGAGATCC	TTTTTTTCTG
	GTCTGGGGCA	TCTTTTCTAG	TTTCCTAGAA	GAACCTAGG	AAAAAAAGAC

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1251	CGCGTAATCT	GCTGCTTGCA	AACAAAAAAA	CCACCGCTAC	CAGCGGTGGT
	GCGCATTAGA	CGACGAACGT	TTGTTTTTTT	GGTGGCGATG	GTCGCCACCA
1301	TTGTTTGCCG	GATCAAGAGC	TACCAACTCT	TTTTCCGAAG	GTAAGTGGCT
	AACAAACGGC	CTAGTTCTCG	ATGGTTGAGA	AAAAGGCTTC	CATTGACCGA
1351	TCAGCAGAGC	GCAGATACCA	AATACTGTCC	TTCTAGTGTA	GCCGTAGTTA
	AGTCGTCTCG	CGTCTATGGT	TTATGACAGG	AAGATCACAT	CGGCATCAAT
1401	GGCCACCACT	TCAAGAACTC	TGTAGCACCG	CCTACATACC	TCGCTCTGCT
	CCGGTGGTGA	AGTTCTTGAG	ACATCGTGGC	GGATGTATGG	AGCGAGACGA
1451	AATCCTGTTA	CCAGTGGCTG	CTGCCAGTGG	CGATAAGTCG	TGTCTTACCG
	TTAGGACAAT	GGTCACCGAC	GACGGTCACC	GCTATTCAGC	ACAGAATGGC
1501	GGTTGGACTC	AAGACGATAG	TTACCGGATA	AGGCGCAGCG	GTCGGGCTGA
	CCAACCTGAG	TTCTGCTATC	AATGGCCTAT	TCCGCGTCGC	CAGCCCGACT
1551	ACGGGGGGTT	CGTGCATACA	GCCAGCTTG	GAGCGAACGA	CCTACACCGA
	TGCCCCCAA	GCACGTATGT	CGGGTCGAAC	CTCGCTTGCT	GGATGTGGCT
1601	ACTGAGATAC	CTACAGCGTG	AGCTATGAGA	AAGCGCCACG	CTTCCCGAAG
	TGACTCTATG	GATGTCGCAC	TCGATACTCT	TTGCGGTGTC	GAAGGGCTTC
1651	GGAGAAAGGC	GGACAGGTAT	CCGGTAAGCG	GCAGGGTCGG	AACAGGAGAG
	CCTCTTTCCG	CCTGTCCATA	GGCCATTGCG	CGTCCCAGCC	TTGTCTCTCTC
1701	CGCACGAGGG	AGCTTCCAGG	GGGAAACGCC	TGGTATCTTT	ATAGTCCTGT
	GCGTGCTCCC	TCGAAGGTCC	CCCTTTGCGG	ACCATAGAAA	TATCAGGACA
1751	CGGGTTTCGC	CACCTCTGAC	TTGAGCGTCG	ATTTTTGTGA	TGCTCGTCAG
	GCCCAAAGCG	GTGGAGACTG	AACTCGCAGC	TAAAAACACT	ACGAGCAGTC
1801	GGGGGCGGAG	CCTATGGAAA	AACGCCAGCA	ACGCGGCCTT	TTTACGGTTC
	CCCCCGCCTC	GGATACCTTT	TTGCGGTCGT	TGCGCCGGAA	AAATGCCAAG
1851	CTGGCCTTTT	GCTGGCCTTT	TGCTCACATG	TTCTTTCCTG	CGTTATCCCC
	GACCGGAAAA	CGACCGGAAA	ACGAGTGTAC	AAGAAAGGAC	GCAATAGGGG
1901	TGATTCTGTG	GATAACCGTA	TTACCGCCTT	TGAGTGAGCT	GATACCGCTC
	ACTAAGACAC	CTATTGGCAT	AATGGCGGAA	ACTACTCGA	CTATGGCGAG
1951	GCCGCAGCCG	AACGACCGAG	CGCAGCGAGT	CAGTGAGCGA	GGAAGCGGAA
	CGGCGTCGGC	TTGCTGGCTC	GCGTCGCTCA	GTCACTCGCT	CCTTCGCCTT
2001	GAGCGCCCAA	TACGCAAACC	GCCTCTCCCC	GCGCGTTGGC	CGATTCAATTA
	CTCGCGGGTT	ATGCGTTTGG	CGGAGAGGGG	CGCGCAACCG	GCTAAGTAAT
2051	ATGCAGCTGG	CACGACAGGT	TTCCCGACTG	GAAAGCGGGC	AGTGAGCGCA
	TACGTCGACC	GTGCTGTCCA	AAGGGCTGAC	CTTTCGCCCC	TCACTCGCGT
2101	ACGCAATTAA	TGTGAGTTAG	CTCACTCATT	AGGCACCCCA	GGCTTTACAC
	TGCGTTAATT	ACACTCAATC	GAGTGAGTAA	TCCGTGGGGT	CCGAAATGTG
2151	TTTATGCTTC	CGGCTCGTAT	GTTGTGTGGA	ATTGTGAGCG	GATAACAATT
	AAATACGAAG	GCCGAGCATA	CAACACACCT	TAACACTCGC	CTATTGTTAA
2201	TCACACAGGA	AACAGCTATG	ACCATGATTA	CGCCAAGCTT	GCATGCAAAT
	AGTGTGTCCT	TTGTCGATAC	TGGTACTAAT	GCGGTTGAA	CGTACGTTTA
2251	TCTATTTCAA	GGAGACAGTC	ATAATGAAAT	ACCTATTGCC	TACGGCAGCC
	AGATAAAGTT	CCTCTGTCAG	TATTACTTTA	TGGATAACGG	ATGCCGTCGG
2301	GCTGGATTGT	TATTACTCGC	GGCCCAGCCG	GCCATGGCCC	AGGTGCAGCT
	CGACCTAACA	ATAATGAGCG	CCGGGTCGGC	CGGTACCGGG	TCCACGTCGA
2351	GCAGGTCGGC	CTCGAGATCA	AACGGGCGGC	CGCAGGTGCG	CCGGTGCCGT
	CGTCCAGCCG	GAGCTCTAGT	TTGCCCGCCG	GCGTCCACGC	GGCCACGGCA
2401	ATCCAGATCC	GCTGGAACCG	CGTGGGGCCG	CAAGCGCTTG	GAGCCACCCG
	TAGGTCTAGG	CGACCTTGGC	GCACCCCGGC	GTTGCGGAAC	CTCGGTGGGC
2451	CAGTTCGAAA	AATAATAAGG	ATCCGAATTC	ACTGGCCGTC	GTTTTACAAC
	GTCAAGCTTT	TTATTATTCC	TAGGCTTAAG	TGACCGGCAG	CAAAATGTTG
2501	GTCGTGACTG	GGAAAACCCT	GGCGTTACCC	AACTTAATCG	CCTTGCAGCA

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	CAGCACTGAC	CCTTTTGGGA	CCGCAATGGG	TTGAATTAGC	GGAACGTCGT
2551	CATCCCCCTT	TCGCCAGCTG	GCGTAATAGC	GAAGAGGCCC	GCACCGATCG
	GTAGGGGGAA	AGCGGTCGAC	CGCATTATCG	CTTCTCCGGG	CGTGGCTAGC
2601	CCCTTCCCAA	CAGTTGCGCA	GCCTGAATGG	CGAATGGCGC	CTGATGCGGT
	GGGAAGGGTT	GTCAACGCGT	CGGACTTACC	GCTTACCGCG	GACTACGCCA
2651	ATTTTCTCCT	TACGCATCTG	TGCGGTATTT	CACACCGCAT	ACGTCAAAGC
	TAAAAGAGGA	ATGCGTAGAC	ACGCCATAAA	GTGTGGCGTA	TGCAGTTTCG
2701	AACCATAGTA	CGCGCCCTGT	AGCGGCGCAT	TAAGCCCGGC	GGGTGTGGTG
	TTGGTATCAT	GCGCGGGACA	TCGCCGCGTA	ATTCGGGCCG	CCCACACCAC
2751	GTTACGCGCA	GCGTGACCGC	TACACTTGCC	AGCGCCCTAG	CCCCCGCTCC
	CAATGCGCGT	CGCACTGGCG	ATGTGAACGG	TCGCGGGATC	GGGGGCGAGG
2801	TTTCGCTTTC	TTCCCTTCCT	TTCTCGCCAC	GTTCGCCGGC	TTTCCCCGTC
	AAAGCGAAAG	AAGGGAAGGA	AAGAGCGGTG	CAAGCGGCCG	AAAGGGGCAG
2851	AAGCTCTAAA	TCGGGGGCTC	CCTTTAGGGT	TCCGATTTAG	TGCTTTACGG
	TTCGAGATTT	AGCCCCGAG	GGAAATCCCA	AGGCTAAATC	ACGAAATGCC
2901	CACCTCGACC	CCAAAAAACT	TGATTTGGGT	GATGGTTCAC	GTAGTGGGCC
	GTGGAGCTGG	GGTTTTTTGA	ACTAAACCCA	CTACCAAGTG	CATCACCCGG
2951	ATCGCCCTGA	TAGACGGTTT	TTCGTCCTTT	GACGTTCGAG	TCCACGTTCT
	TAGCGGGACT	ATCTGCCAAA	AAGCAGGAAA	CTGCAAGCTC	AGGTGCAAGA
3001	TTAATAGTGG	ACTCTTGTTT	CAAACCTGAA	CAATACTCAA	CCCTATCTCG
	AATTATCACC	TGAGAACAAG	GTTTGACCTT	GTTATGAGTT	GGGATAGAGC
3051	GGCTATTCTT	TTGATTTATA	AGGGATTTTG	CCGATTTTCG	CCTATTGGTT
	CCGATAAGAA	AACTAAATAT	TCCCTAAAAC	GGCTAAAGCC	GGATAACCAA
3101	AAAAAATGAG	CTGATTTAAC	AAAAATTTAA	CGCGAATTTT	AACAAAATAT
	TTTTTTACTC	GACTAAATTG	TTTTTAAATT	GCGCTTAAAA	TTGTTTTATA
3151	TAACGTTTAC	AATTTTATGG	TGCAGTCTCA	GTACAATCTG	CTCTGATGCC
	ATTGCAAATG	TTAAAATACC	ACGTCAGAGT	CATGTTAGAC	GAGACTACGG
3201	GCATAGTTAA	GCCAGCCCCG	ACACCCGCCA	ACACCCGCTG	ACGCGCCCTG
	CGTATCAATT	CGGTCGGGGC	TGTGGGCGGT	TGTGGGCGAC	TGCGCGGGAC
3251	ACGGGCTTGT	CTGCTCCCGG	CATCCGCTTA	CAGACAAGCT	GTGACCGTCT
	TGCCCCGAACA	GACGAGGGCC	GTAGGCGAAT	GTCTGTTTCA	CACTGGCAGA
3301	CCGGGAGCTG	CATGTGTCAG	AGGTTTTTAC	CGTCATCACC	GAAACGCGCG
	GGCCCTCGAC	GTACACAGTC	TCCAAAAGTG	GCAGTAGTGG	CTTTGCGCGC
3351	A				
	T				

Fig. 8b

cDNA primers

VLK-c	CTGGATGGTGGGAAGATGGA (SEQ ID No:113)
VLL-c	TCAGAGGAAGGAAACAGGGT (SEQ ID No:114)
IgG1-c	CTTACAACCACAATCCCTGGGCACAATTTT (SEQ ID No:115)
IgG2a-c	CTTTGTGGGCCCTCTGGGCTCAAT (SEQ ID No:116)
IgG2b	TGAAATGGGCCCCGCTGGGCTCAAG (SEQ ID No:117)
IgG3-c	GGGCTTGGGTATTCTAGGCTCGAT (SEQ ID No:118)

VH forward primers without restriction sites

M-VH1	GAGGTGCAGCTTCAGGAGTCAGG (SEQ ID No:119)
M-VH2	CAGGTGCAGCTGAAGGAGTCAGG (SEQ ID No:120)
M-VH3	GAGGTCCAGCTGCAACAGTCTGG (SEQ ID No:121)
M-VH4	GAGGTTTCAGCTGCAGCAGTCTGG (SEQ ID No:122)
M-VH5	CAGGTCCAACCTGCAGCAGCCTGG (SEQ ID No:123)
M-VH6	CAGGTTTCAGCTGCAGCAGTCTGG (SEQ ID No:124)
M-VH7	GAGGTGAAGCTGGTGGAGTCTGG (SEQ ID No:125)
M-VH8	GAGGTGAAGCTGGTGGAACTCTGG (SEQ ID No:126)
M-VH9	GAGGTTTCAGCTTCAGCAGTCTGG (SEQ ID No:127)

VH backward primers without restriction sites

M-JH1	TGAGGAGACGGTGACCGTGGTCCC (SEQ ID No:128)
M-JH2	TGAGGAGACTGTGAGAGTGGTGCC (SEQ ID No:129)
M-JH3	TGCAGAGACAGTGACCAGAGTCCC (SEQ ID No:130)
M-JH4	TGAGGAGACGGTGACTGAGGTTCC (SEQ ID No:131)

VL forward primer without restriction sites

M-VK1	GACATTGTGATGACACAGTCTCC (SEQ ID No:132)
M-VK2	GATGTTGTGATGACCCAACTCC (SEQ ID No:133)
M-VK3	GATATCCAGATGACACAGACTCC (SEQ ID No:134)
M-VK4	CAAATTGTTCTCACCCAGTCTCC (SEQ ID No:135)
M-VL1	CAGGCTGTTGTGACTCAGGAATC (SEQ ID No:136)

VL backward primers without restriction sites

M-JK1	TTTGATTTCCAGCTTGGTGCCTCC (SEQ ID No:137)
M-JK2	TTTTATTTCCAGCTTGGTCCCCC (SEQ ID No:138)
M-JK3	TTTCAGCTCCAGCTTGGTCCCAGC (SEQ ID No:139)
M-JL1	ACCTAGGACAGTGACCTTGGTTCC (SEQ ID No:140)

VH forward primers with restriction sites

MVH1 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGCAGCTTCAGGAGTCAGG (SEQ ID No:141)
MVH2 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGCAGCTGAAGGAGTCAGG (SEQ ID No:142)
MVH3 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTCCAGCTGCAACAGTCTGG (SEQ ID No:143)
MVH4 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTTCAGCTGCAGCAGTCTGG (SEQ ID No:144)
MVH5 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTCCAACTGCAGCAGCCTGG (SEQ ID No:145)
MVH6 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTTCAGCTGCAGCAGTCTGG (SEQ ID No:146)
MVH7 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGAAGCTGGTGGAGTCTGG (SEQ ID No:147)
MVH8 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTGAAGCTGGTGGAACTCTGG (SEQ ID No:148)
MVH9 Sfil	GTCCTCGCAACTGCGGCCCAGCCGGCCATGGCCGAGGTTCAGCTTCAGCAGTCTGG (SEQ ID No:149)

VH backward primers with restriction sites

MJH1 Sall	GAGTCATTCTCGTGTGACACGGTGACCGTGGTCCC (SEQ ID No:150)
MJH2 Sall	GAGTCATTCTCGTGTGACACTGTGAGAGTGGTGCC (SEQ ID No:151)
MJH3 Sall	GAGTCATTCTCGTGTGACACAGTGACCAGAGTCCC (SEQ ID No:152)
MJH4 Sall	GAGTCATTCTCGTGTGACACGGTGACTGAGGTTCC (SEQ ID No:153)

VL forward primers with restriction sites

MVK1 ApaL1	TGAGCACACAGTGCACTCGACATTGTGATGACACAGTCTCC (SEQ ID No:154)
MVK2 ApaL1	TGAGCACACAGTGCACTCGATGTTGTGATGACCCAACTCC (SEQ ID No:155)
MVK3 ApaL1	TGAGCACACAGTGCACTCGATATCCAGATGACACAGACTCC (SEQ ID No:156)
MVK4 ApaL1	TGAGCACACAGTGCACTCCAAATTGTTCTCACCCAGTCTCC (SEQ ID No:157)
MVL1 ApaL1	TGAGCACACAGTGCACTCCAGGCTGTTGTGACTCAGGAATC (SEQ ID No:158)

VL backward primers with restriction sites

M-JK1 Not1	GAGTCATTCTCGACTTGCGGCCGCTTTGATTTCCAGCTTGGTGCCTCC (SEQ ID No:159)
M-JK2 Not1	GAGTCATTCTCGACTTGCGGCCGCTTTTATTTCCAGCTTGGTCCCCC (SEQ ID No:160)
M-JK3 Not1	GAGTCATTCTCGACTTGCGGCCGCTTTCCAGCTCCAGCTTGGTCCCAGC (SEQ ID No:161)
M-JL1 Not1	GAGTCATTCTCGACTTGCGGCCGACCTAGGACAGTGACCTTGGTTCC (SEQ ID No:162)

Fig. 9